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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 03-JUN-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (416) 595-11 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 898 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
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APPLICANT: Yang, Yan-Ping
APPLICANT: Klein, Michel H
APPLICANT: Klein, Michel H
TITLE OF INVENTION: LACTOFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                         344 SLTGKLSYYDNPNQQTAQGKYIKSQFDTTKKVNETDVYQIDAKINGNRFVGTAKSLVNEN 403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 y Match 4.5%; Score 178; DB 2; I
Local Similarity 18.6%; Pred. No. 4.9e-06;
hes 171; Conservative 118; Mismatches 305;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC OPERATING SYSTEM:
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STATE: Ontario
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                                       FQARVFSSND-----IGERIELSDGTYTLTELNSPAGYSIAEPITFKVEAGKVYTIIDG
                                                                                           TETAPFIKELFSKKANPNNP--NPN--
--AGKFLSNDNASYVVFGGKRDKTDKPVATKTVYFSAGFE--KPSTSFVDNETIGRIINS 498
                                                                                                                                                                                                                                                                                                                                                                                    GSSLTQKNADTPNDKDRIPKPMPILFYHG--ENASSQLPSAGKFNYTGNWLYLSDVKKRP 292
                                                                                                                                       L------VPTKPPTPGDPPMPPNQPQTTSVLIRKYAIGDYSKLLEGATLQLTGDNVNS
                                                                                                                                                                                                                                                                                      ALSASDDRVGVYLNASGKSN------EGDVVSAAHIYLNGFQYKHTPATYQVDFDTN 343
                                                                                                                                                                                                                                                                                                                                     AIRVTQEAVWYYSDNAPISNPDESFKRESESNLVSTSQLSL-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EGSRSYQVYCFNLKKAFPLGSDSSVKK--WYKKHDGISTKFEDYAMSP------RIT 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PDYSNDLVQQWQGKLYVGIDAHRPDGIGTGKNLRQPITANDIKPLYFNKFPALSDLHLDS 175
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6th Floor, 330 University Avenue
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(416) 595-1163
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SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version
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US-08-867-941-11
Sequence 11, Application US/08867941
Patent No. 5977337
GENERAL INFORMATION:

APPLICANT:

Loosmore,

Sheena M

RESULT

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RESULT 3
US-09-074-658-70
US-09-074-658-70; Sequence 70, Application US/09074658; Patent No. 6184371
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                                                                           APPLICATION NUMBER: US/09/074,658
APPLICATION UMBER: US/09/074,658
APPLICATION: 08-WAY-1998
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Stewart, Michael I
REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 1038-795
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
(416) 595-1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                  TELEFAX: (416) 595-1163 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
              SEQUENCE CHARACTERISTICS:
LENGTH: 905 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Klein, Michel H
TITLE OF INVENTION: LACTOFERRIN RECEPTOR GENES OF MORAXELLA
NUMBER OF SEQUENCES: 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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ZIP: M5G 1R7
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Yang, Yan-Ping
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Sim a McBurney
י פוחסד, 330 University Avenue
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                                                                                                                                                                                                                                                              Version
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; STRANDEDNESS: single ; TOPOLOGY: linear US-09-074-658-70
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Best Local Similarity
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862 IDVKGRFFGTNGEELA
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                                                                                                                                                                                                                                                                                                                                                                                                                                               548 KNPAYFGQHDKFYFNGNYYDLSAKEANKLGVSQDTSTNKSILAKYPDA----KVSTDNKV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              334 YKHTPATYSVDFDQNTLKGKLSYYDNPNKQTADGRYIRSQFDTDKKVNEADVYEIDAKIN
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                                TDSEGYKVKVNSQEVA 696
                                                                                                        KDFHFEIELKNNKQELLSQTVKTDKTNLEFKDGKATINLKHGESLTLQGLPEGYSYLVKE
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                                                                       GDIVFDI - - KNGKIDGTGFTAKADVPNYREEVG - -
                                                                                                                                                GKVYHAGN----IRPEFENKYLPINEPTHEKTFALDGKNKAKFDVNFDTNSLTGKLNDER
                                                                                                                                                                                 GTQWHPEDLVDIIRMEDKKEVIPV---THNLTL-----
                                                                                                                                                                                                                       DSLTASDDSQDDNTHGDDDL-----IASDDSQDDDTDGDDDSDDL--GDGADDDAA
                                                                                                                                                                                                                                                        DSAELDKDKLKD-YHGFGDMNDSTLAVAKILVEYAQDSNPPQLTDLDFFIPNNNKYQSLI
                                                                                                                                                                                                                                                                                             FTYNGLWAGYLTQKKDKGYSKDEDTIKQKGLKD-----YILTKDFIPQDDDDDD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QNYA-----KFYY-----AK--NKNGSSQVVYCFNADLKSPPDSEDGGKTMTPDFT 449
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LEGATLQLTGDNVNSFQARVFSSNDI------GERIELSDGTYTLTELNSPAGYSIAEP 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GDKYNKGYQNLLSGGL------VPTKPPTPGDPPMPPNQPQTTSVLIRKYAIGDYSKL 306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KNKKPGDDYQNIRFGYMELRELDLNKKGADNQSDKNRAIIFTTPTLFYHGENASTHLP--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STK--FEDYA-----MSPRIT-----GDELNQKLRAVMYNG-----HPQNANGIMEGL 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YFDKFPKISDL--HLENSE----HVFDAKKANNIKIYGYGALSSPAKNPTYMNYQQEQNI 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative 102; Mismatches 264; Indels 266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.5%; Score 176; DB 4; 20.6%; Pred. No. 7.2e-06;
877
                                                                                                                                                                                                                                                                                                                                                                                                           ----GEVKYTHIAGRDLFKYTVKPRDTDPDTFLK----
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US-09-074-658-15

Sequence 15, Application US/09074658 Patent No. 6184371 GENERAL INFORMATION:

APPLICANT: APPLICANT:

Run-Pan

Loosmore, 5

Sheena M

RESULT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 24,973
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (416) 595-1155
TELEFAX: (416) 595-1163
INFORMATION FOR SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Quijun Wang
APPLICANT: Yang, Yan-Ping
APPLICANT: Klein, Michel H
TITLE OF INVENTION: LACTOFERRIN RECEPTOR GENES OF MORAXELLA
NUMBER OF SEQUENCES: 78
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sim & McBurney
STREET: 6th Floor, 330 University Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: Canada
ZIP: M5G 1R7
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/074,658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 2432 amino acids
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ATTORNEY/AGENT INFORMATION:
NAME: Stewart, Michael I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 6th Flo
CITY: Toronto
STATE: Ontario
                                                                                        481 KPSTSFVDNETIGGIIDRKGLNNHINEDETIPSDDSYYGYTWGKPEKQFTKKVSSSTQVV 540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             179
541 PAYFGQHDKFYF----NGNYYDLSASRVDKLAPADAVKANQSIKEKYPNATLNKDNQVTA 596
                                          407 ----NYAKFYYAKNKNGSSQVVYCFNADLKSPPDSEDGGKTMTPDFTTGEV-----
                                                                                                                                                                                    305 KLLEGATLQLTGDNVNSFQARVFSSNDI------GERIELSDGTYTLTELNSPAGYSIA 357
                                                                                                                                                                                                                                                                                  380 VYQIDAKINGNRFVGTAKSLVNEKTQTAPFIKELFSKKANPNNPNPN------S
                                                                                                                                                                                                                                                                                                                                 262 -YQ-----NLLSG---GLVPTKPPTPG-----DPPMPPNQPQTTSVLIRKYAIGDYS 304
                                                                                                                                                                                                                                                                                                                                                                                329
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         278 YTGNWLYLSDVKKRPALSASDERVGVYLNASGKAN-----EGDVVSAAHIYLNGFQ 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         220 VDPYENIRFGYLELQGSSLTQKNADNQNEQDRIPKPMPILFYHG--ENASSQLPSAGKFN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      142 MSP-----RITGDELNQK------LRAVMYNGHPQNANGIMEGLEPLN 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     161 YFNKFPALSDLHLDSERHR-FDPQKINTIKVYGYGNLTTPSNNNTHINHQQADNKKNNKP 219
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TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                84 YYKQFRVAHDLRVNLEGSRSYQVYCFNLKKAFPLG--SDSSVKKWYKKHDGISTKFEDYA 141 | : | : | : | : | : | : |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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                                                                                                                                     EPITFKVEAGKVYTTIDGKQIENP-NKETVEPYSVEAY-----NDFEEFSVLTTQ-- 406
                                                                                                                                                                                                                                                                                                                                                                             YKH-----TPATYQVDFDTNSLTGKLSYYDNPNQ--QNNKGEYLKSQFDTTKKVNETD 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ontario
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	NUMBER	٠.	
	TRATION NUMBER: 39,300	٠. ٠	
	NAME: McGowan, Malcolm K.		
	FILING DATE:	٠.	
	APPLICATION NUMBER:	 .	
	PLICATION DATA:	٠	
	APPLICATION NUMBER: WO PCT/SEVI/00/0/		
	A.	•••	
	: 21-AUG-1992		
	BER. IIS		
	CLASSIFICATION: 435	٠.	
	2-MAY-1995	٠	
	APPLICATION NUMBER: US/08/447,031A		
	ARE: PatentIn Release #	٠	
	SYSTEM: PC-DOS/MS-DOS	•. •	
	COMPUTER: IBM PC compatible	·. ·	
	READABLE FORM:		
	22313-1404	••	
	COUNTRY: United States		
	CITY: ALEXANORIA		
	T: P.O. Box 1404	٠.	
	SSEE: Burns,		
	NOMBER OF VECCENCES: 0		
	OF INVENTION: ITS PREPARA	٠	
	OF INVENTION: A CO	٠. ،	
	APPLICANT: SIGNAS, CHLISTER APPLICANT: SWITALSKI, Lech		
	: PATTI, Jose		
	: LINDBER	••	
		٠	
	APPLICANT: GUSS, Bengt		
	ORMATION:	٠.	
	Sequence 2, Apprication 05/0044/031A Patent No. 5851794		
	031A-2	RE	
	STd	Db	
	728 KINGYLALIVIAGISLGIW 746	Qy	
ттн эт	854 KGQFFGTNGEELAGQLHHDKGDGINDTAEKAGAVFGAVKDKMSKS11K1QTFSVH1MTTH	뫄	
72		Qy	
٥	810 VEDLKNGKLUGTGETAKADVENIKEEVGNNQGGGELENIAD	D	
D J		! . £	
TDS 683	624 HFETELKNNKOELLSOTVKTDKTNLEFKDGKATINLKHGESLTLOGLPEGYSYLVKETDS	Q	
GDI 809	754 YHAGNIRPEFENKYLPINEPTHEKTFALDGKNKAKFEVDFNTNSLTGKLNDERGDI	뫄	
KDF 623	582 WHPEDLVDIIRMEDKKEVIPVTHNLTLRTTVTGLAGDRTKDF	Qy	
ĠKV 753	705 TASDDSQDDNTHGDDDLIASDDSQDDDADGDDDSDDLGDGADDDAAGKV	Db	
GTQ 581	522 AELDKDKLKDYHGFGDMNDSTLAVAKILVEYAQDSNPPQLTDLDFFIPNNNKYQSLIGTQ	Qy	
DDL 704	657 TYNGLWAGYLTQKKDKGYSDNAETIKEKGHPGYLLTENFTPEDDDDDL	В	
-DS 521 	486 IEKGYREKGQAIEYSGLTETQLRAATQLAIYYFT	Qy	
GKF 656	597 IVLQEAKDNKPYTAIRAKSYQHISFGETLYNDANQTPTRSYFVQGGRADTSTTLPQAG	рь	
- KV 48	454KYTHIA-GRDLFKYTVKPRDTDPDTELKHIKKV	Qy	

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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (703) 836-20 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 1183 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      296
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           398 PYTFDKDKEYPFTMKDTDNQGYFTTIENAKAIEKTKDVSAQKVWEGTQKVKP-----TI
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                                                                                                                                                             LPKYDEGKKI-----EYTVT---EDHVKDYTTDINGTTITNKYTPGETSATVTKNW
TKVKGYTTHVDNNDMGNLIVTNKYTPETTSISGEKVWDDKDNQDGKRPEKVSVNLLANG
                                                                                                                    -DKTNLE-----FKDGKAT----INLKHGESLTLQGLPE-----GYSYLVKE- 680
                                                                                                                                                                                                                                                                               HGFGDMNDSTLAVAKILVEYAQDSNPPQLTDLDFFIPNNNKYQSL---IGTQWHPEDLVD
                                                                              DDNNNQDGKRPTEIKVELYQDGKATGKTAILNESNNWTHTWTGLDEKAKGQQVKYTVEEL
                                                                                                                                                                                                   IIRMEDKKEVIPVTHNLTLRKTVTGLAGDRTKDFHFEIE-----LKNNKQELLSQTVKT- 643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VEPYSVEAYNDFEEFSVLTTQNYAKFYY-----AKNKNGSSQVVYCFNADLKSPPDSEDG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YTLTELN-----SPAGYSIAE------PI-TFKVEAGKVYTIIDGKQIENPNKEI
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                                       TDSEGYKVKVNSQEVANATVSK-----TGITSDETLAFENNKEPVVPTGVDQKI--NG
                                                                                                                                                                                                                                        ----DKDN------QDGKRPEKVSVN-LLADGEKVKTLDVTSETNWKYE-FKD
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                                                                                                                                                                                                                                                                                                                                                            -----GYREKGQAIEYSGLTETQLRAATQ------LAIYYFTDSAELDKDKLKDY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ----DKGDKY-----NKGYQNLLSGGL------VPTKPPTPGDPPMPPNQPQTTSV 293
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1112 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,402
FILING DATE: 16-SEP-1996
FILING DATE: 16-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                        263 -QNLLS-----
                                                                                                                                                                                                                                                                                                                       218 -ISQDIPKIIYDSANSPLAIGKYHAENHQLIYTFTDYIAGLDKVQLSAELSLFLENKEVL 276
                                                                                                                                                                                                                                                                                                                                                                                                        167 TEVSSGAADFYRNHAAYFKMSFELKQKDKSETINPGDTFVLQLDRRLNPKG------ 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            y Match 3.7%; Score 145; DB 2;
Local Similarity 19.8%; Pred. No. 0.0035;
hes 190; Conservative 121; Mismatches 327
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STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                           AVWYYSDNAPISN-PDESFK-----RESESNLVSTSQLSLMRQALKQLIDPNLATKMP 237
                                                               SSYGVDVTKLTLRTDITAGLGNGFQMTKRQRIDFGNNIQNKAFIIKVTGKTDQSGKPLVV 454
                                                                                                                                                 FVWYVYVNPNRTNIPYATMNLWGFGRARSNTSDLENDANTSSAELGEIQVYEVPEGEKLP
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                                                                                                                                                                                                                                                                             -----MSPRITGDELNQK--LRAVMYNGHPQNANGIMEGL-----EPLNAIRVTQE 185
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FISCHETTI, Vincent A.
FISCHETTI, Vincent A.
FIBRONECTIN AND FIBRINGEN BINDING
VENTION: FIBRONECTIN AND FIBRINGEN BINDING
VENTION: FIBRONECTIN AND FIBRINGEN BINDING
VENTION: FIBRONECTIN AND FIBRINGEN BINDING
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-GG-----LVPTKPPTPGDPPMPPNQPQTTSVLIRKYA 299
                                                                                                       ----FQLSIFESEDKGDK-YNKGY------
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                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                      SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                   PRIOR APPLICATION DATA:
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                                  FILING DATE: 2: CLASSIFICATION:
APPLICATION NUMBER:
                                                                     APPLICATION NUMBER:
                                                                                                                           OPERATING SYSTEM: MS-DOS Version
                                                                                                                                            COMPUTER: Hewlett-Packard Vectra
                                                                                                                                                                                                        ZIP: 63043
                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                              CITY: St.
                                                                                                                                                                                                                                                                              STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R-ETEFVFEQNNFNLEFGNAEIKGQSGSKIIDEDTLTSFKGKKIWKNDTAENRPQAIQVQ 767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KI 729
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       II -- NTKREVITQQGPKLEIEETLPLESGASG--GTTTVEDSRPVDTLSGLSSEQGQSGD
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                                                                                                                                                                                                                                           Missouri
                                                                                                                                                                                                                                                                            929 Fee Fee Road,
                                                                                                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                     Wise, Kim 5.
McIntosh, Mark A.
                                                                                                                                                                                                                                                              Louis
                                                                                                                                                                                                                                                                                                   Grace J.
                                                    28-AUG-1996
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                                                                                                                                                                                                                                                                                                                                                                   DNA Sequences Coding for Mycoplasma
Hyopneumoniae Surface Antigens,
Corresponding Proteins and Use in
                                                                                                                                                                                                                                                                                                                                                         Vaccines
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08/373,957
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; PUBLICATION DATE: RELEVANT RESIDUES IN SEQ US-08-703-947-2
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DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PUBLICATION INFORMATION:
AUTHORS: Mary F. Kim,
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LIBRARY: Genom
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627 residues of the P65 lipoprotein, deriv
from the nucleic acid sequence
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US-08-480-604A-10
Sequence 10, Application US/08480604A
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GENERAL INFORMATION:
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                                                   APPLICANT: KINK, JOHN A.
APPLICANT: THALLEY, BRUC
APPLICANT: PADHYE, NISH!
   APPLICANT: FIRCA, APPLICANT: STAFFORITIFLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NFLAEKGMLLKAEVNKKIEEINTNLDNLIKELKALNPKLSINLVGYKLPNSG------
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                                   THALLEY, BRUCE S. PADHYE, NISHA V. FIRCA, JOSEPH R.
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 STAFFORD, DOUGLAS C.
VENTION: VACCINE AND ANTITOXIN FOR TREATMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ----NKPTYSKVI-----
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US-08-480-604A-10
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                                                                                                                                                                                                                                                                                                                                           Matches 154;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 2366 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION UNMEER: US 08/329,154
FILING DATE: 25-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: PR
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                               1738 IRYVW-----SNDGNDF-----ILMSTSEENKVSQVKIRFVNVFKDKTLANKLS--
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US 08/161,907
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MEDIUM TYPE: Floppy disk
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   285
                                                                                                                                                   184 QEAVWYYSDNAPISNPDESFKRESESNLVSTSQLSLMRQALKQLI----DPNLATKMPKQ
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FTLING DATE: 14-APR-1995
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STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/985,321 FILING DATE: 04-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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 PNQPQTTSVLIRKYAIGDYSKLLEGATLQLTGDNVNSFQARVFSSNDIGERIELSDGTYT 344
                                                                          VPDDFQLSIFESEDKGD-----
                                                                                                                                                                                                                                                                 PYFIKFNTLETNYTLYVGNRQNMIVEPNYDLDDSGDISSTVINFSQKYLYGIDSCVNK--
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220 MONTGOMERY STREET, SUITE
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                                                                                                                                                                                          ----VVISPNIYTDEINITPVYETNNTYPEVIVLDANYINEKIN-VNINDLS
                                       -FNFSDKQDVPVSEIILSFTPSY---YEDGLIGYDLGLV---
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31-OCT-1989
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                                                                                                                                                                                                                                                                                                                                           86; Mismatches 249;
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284

1781 239 1737 DB 1; Length 2366;

Indels 296;

Gaps

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RESULT 9
US-08-405-496A-10
; Sequence 10, Application US/08405496A
; Patent No. 5919665
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                                  PRIOR APPLICATION DATA:
APPLICATION WIMBER: US 08/329,154
FILING DATE: 25-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/161,907
FILING DATE: 02-DEC-1993
                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2304
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                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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                    PRIOR APPLICATION DATA:
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                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                      APPLICATION NUMBER: US/0 FILING DATE: 16-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: MEDLEN & CARROLL, LI
STREET: 220 MONTGOMERY STREET,
CITY: SAN FRANCISCO
                                                                                                                                                                                                                                                                                                                                                                      STATE:
APPLICATION NUMBER:
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US 07/985,321
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Best Local Similarity
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TELEFAX: (415) 397-8338
INFORMATION FOR SEQ ID NO:
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LENGTH: 2366 amino acids
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FILING DATE: 31-OCT-1989
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           240
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                                                                                                                                                                                                                                                                         ------LDKDKLKDYHGFGDMN------DSTLAVAKILVE----
                                                                                                                                                                                                                                                                                                                                                                                                             DGVMQKGFVSINDNKHYFDDSGVMKVGYTEIDGKHFYFAENGEMQIGVFNTEDGFKY---
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                                               FGETYTIETGW----IYDMENESDKYYFNPET----KKACKGINLIDDIKYYFDEKGIM
                                                                                                                                       VQNIDDNYFYIDDNGIVQIGVFDTSDGYKYFAPANTVNDNIYGQAVEYSGLVRVGEDVYY 2200
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                                                                                       ----IGTQWHPEDLVDIIRMEDKKEVIPVTHNLTLRKTVTGLAGDRTKDFHFE-----
                                                                                                                                                                                 -----YAQDSNPPQLTDLD-----FFIPNNN------KYQSL------
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-IELKNNKQELLSQTVKTDKTNLEFKDGKATINLK---HGESLTLQ----GLPEGYS
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US-08-867-941-15
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Best Local Similarity 19.7:
Conservative
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Sequence 15, App....
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, '
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/867,941
FILING DATE: 03-JUN-1997
CLASSIFICATION: 435
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APPLICANT:
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REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
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NAME: Stewart, Michael I
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                                                                                                                                                                                                                         142
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ZIP: M5G 1R7
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                                    YKH-----TPATYQVDFDTNSLTGKLSYYDNPNQ--QNNKGEYLKSQFDTTKKVNETD 379
                                                                                                                                                                                                                     MSP-----RITGDELNQK------LRAVMYNGHPQNANGIMEGLEPLN 178
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                                                                                                                                                                                                                                                                                             YYKQFRVAHDLRVNLEGSRSYQVYCFNLKKAFPLG--SDSSVKKWYKKHDGISTKFEDYA 141
 -- yy--
                                                                       LKQLIDPNLATKMPKQVPDDF-----QLSIFESEDKGDKYNKG------
                                                                                                           YTGNWLYLSDVKKRPALSASDERVGVYLNASGKAN-----EGDVVSAAHIYLNGFQ 328
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Yang, Yan-Ping
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---NLLSG---GLVPTKPPTPG-----DPPMPPNQPQTTSVLIRKYAIGDYS 304
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19.7%; Pred. No. 0.0093;
ative 91; Mismatches 283; Indels 2
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                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
               FILING DATE: 23-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: LOGATY, KACTORY
REGISTRATION NUMBER: 36,317
                                                                                                   CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/026,939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Weiser, Jeffrey N. APPLICANT: Plaut, Andrew G.
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                                                                                                                                                                   APPLICATION NUMBER: US/0 FILING DATE: 29-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Philadelphia
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
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                                                                                                                                                                                                                                Release #1.0, Version #1.30
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7600-4U1
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RESULT 12
PCT-US95-10661A-2
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                                                                               1158 AKTKENLANTLEKLLSADAVIASENQTINGYV 1189
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LENGTH: 1861 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19 LSKNSKRFTVTLVGVFLMIFALVTSMVGAKTVFGLVESSTPNAINPDSSSEYRWYGYESY 78
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                                                                                                                                                                                                                                                                                                                                                                                                          SLEERGTILDVSQIASKKSEINSLSAPKVETLLTSTNKESDFSKVKDY----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IKGRFTGKLF-----GSKDGKNYAIY--NLKKPLFDTLSAATVENLTLKDVNISGKT 758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LAKKAEREVTNFTSFSNLVQAINNNLNG---TYYLAASLNANEVELEN-----GASSY 708
                                                                                                                                                            ----GLIYTPNQFLQNHSSIVNEVLPDLKAVDYQSEAIRNTLGISSGVSLTELYLEEQF
                                                                                                                                                                                                  INLKHGESLTLQGLPEGYSYLVKETDSEGYKVKVNSQEVANATVSKTGITSDETL---AF
                                                                                                                                                                                                                                                                                   EVIPVTHNLTLRKTVTGLAGDRTKDFHFEIELKNNKQELLSQTVKTDKTNL-EFKDGKAT 656
                                                                                                                                                                                                                                                                                                                         -----QASRALAYKNIEKLLPFYNKATIVKYGNLVKEDSTLYEKEILSAVMMKD--
                                                                                                                                                                                                                                                                                                                                                               VAKILVEYAQDSNPPQLTDLDFFIPNNN-----KYQSLI--GTQWHPEDLVDIIRMEDKK 597
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                                                                                                                       ----GVDQKINGYL 733
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; MOLECULE TYPE:
PCT-US95-10661A-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 163;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 08/296,791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS LENGTH: 1394 amino ac
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TITLE OF INVENTION: Haemophilus Adherence and Penetration Protein
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: FP-59941/RFT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION: NAME: Trecartin, Richard
295 SLYTRAGNGVYTISGNDNGQG-SITQKSGIPSEIKITLANMSLPLKEKDKVHNPRYDGPN 353
                                                                                                                                                                                                                                                           145 RLHKFYTEAAPIDMTSNMNGSTYSD-
                                                                                                                                                                                                                                                                                                                                                                                     148 GDELNQKLRAVMYNGHPQNANGIMEGLEPLNAIRVTQEAVW---YYSDNA-----PISNP 199
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CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                90 ----NOYIVSVAHNVGYTDVDFGAEGNNP-DQHRFTYKIVKRNNYKKDNLHPYEDDYHNP 144
                                                                                                                                                                                                                                                                                                                                                                                                                                    49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91 -- AHDLRV-NLEGSRSYQVYCFNLKKAFPLGSDSSVKKWYKKHDGISTKFEDYAMSPRIT 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            48 KTVF------GLVESSTPNAINPDSSSEYRWYGYESYVRGHPYYKQFRV-----
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                                                                                  PIAGSKGDSGSPMFIYDAEKQKWLINGILREGNPFEGKENGFQLVRKSYFDEIFERDLHT 294
                                                                                                                                                                      DKGDQVAGAYHYLTAGNTHNQRGAGNG------YSYLGGDVRKAGEYGPL
                                                                                                                                                                                                              DKGDKYNKGYQNLLSGGLVPTKPPTPGDPPMPPNQPQTTSVLIRKYAIGDYSKLLEGATL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1394 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative 117; Mismatches 281;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.2%; Score 128; DB 19.0%; Pred. No. 0.12;
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US-08-946-475-2
; Sequence 2, Application US/08946475
; Patent No. 6013505
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/027,973
FILING DATE: 08-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Gimmi, Edward R
REGISTRATION NUMBER: 38,891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Palmer, Leslie
TITLE OF INVENTION: TOPOISOMERASE
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Gwynn, Michael
APPLICANT: Kallender, Howard
APPLICANT: Palmer, Leslie
                                                                                                                                                       REGISTRATION NUMBER: 38,891
REFERENCE/DOCKET NUMBER: P50560
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             717
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OPERATING SYSTEM: DOS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/946,475 FILING DATE: 08-OCT-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 709 Swedeland CITY: King of Prussia
                                                                                                                   TELEFAX:
                                                                                                                                                   TELEPHONE:
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; TYPE: amino acid
; STRANDEDNESS: Single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-946-475-2
US-08-946-475-9
US-08-946-475-9
; Sequence 9, Application US/08946475
; Patent No. 6013505
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Best Local Similarity
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                                                                                                                                                          661 HGESLTLQGLP 671
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 353 MRTPDDMKSF--LTKDQYRLYKLIWERFVASQMA------PAILD---TVSLDITQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            295 IGLITYMRTDSTRISDTAKAEAKQYITDKY--GESYTSKRKASGKQGDQDAHEAIRPSST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    253 ANPFTTS------TLQQEAARKLNFKARKTMMVAQQLYEGIDLKKQGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          286 NQPQTTSVLIRKYAIGDYSKLLEGATLQLTGDNVNSFQAR--VFSSNDIGERIEL-SDGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 213 STSQLSLMRQALKQLIDPNLATKMPKQV------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             170 IMEGLEPLNAIRVTQEAVWYYS------DNAPISNP-----DESFK--RESESNLV 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 KKYKVIASMGHVRDLPRSQMGVDTEDNYEPKYITIRGKGPVVKELKK-----HAKKAKN 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                        -GEDCEVCGSP 585
                                                                                                                                                                                                                                                                                                                                                         --MTLYVETKDDSDSEKENKLPKLEQ----GD----KVTATQIEPAQHYTQPPPRYTE
                                                                                                                                                                                                    VEFTVNMETLLDKIAEGDITWRKVIDGFFSSFKQDV--ERAEEEMEKIEIKDEPA----
                                                                                                                                                                                                                                        VTHNLTLRKTVTGLA-GDRTKDFHFEIELKNNKQELLSQTVKTDKTNLEFKDGKATINLK 660
                                                                                                                                                                                                                                                                                                                                                                                                                                       GDIK----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VEAYNDFEEFSYLTTQNYAKFYYAKNKNGSSQVVYCFNADLKSPPDSEDGGKTMTPDFTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                YTL-----TELNSPAGYSIAEPITFKVEAGKVYTI---IDGKQIENPNKEIVEPYS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WTIEGEFRYKKSKFNAKFLHYKNKPFKLKTKKDVEKITAALDGDQFEITNVTKKEKTRNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QLSIFESEDKGDKYNK---GYQN-------LLSGGLVPTKPPTPGDPPMPP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DAQQ----ARRILDRLVGYNISPVLWKKVKKGLSAGRVQSVALRLVIDRENEIRNFKPEEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VFLASDP----DREGEAIAWHLSKILELEDSKENRVVFNEITKDAVKESFKNPREIEMNLV 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        691 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                     3.2%; Score 125.5; DB 3;
19.8%; Pred. No. 0.066;
ative 89; Mismatches 228;
                                                                                                                                                                                                                                                                                                                                                                                                                                     ---FRANGQTIKFKGF----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----PDDF 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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GENERAL INFORMATION:
APPLICANT: Gaynn, Michael
APPLICANT: Kallender, Howard
APPLICANT: Palmer, Leslie
TITLE OF INVENTION: TOPOISOMERASE

CORRESPONDENCE NUMBER OF SEQUENCES:

ADDRESS

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SmithKline Beecham Corporation

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-946-475-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity Matches 133; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FASTSEQ for Windows Version CURRENT APPLICATION NUMBER: US/08/946,475 FILING DATE: 08-OCT-1997 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: P5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 08-OCT-1996
ATTORNEY/AGENT INFORMATION:
451 GEVKYTHIAGRDLFKYTVKPRDTDPDTFLKHIKKVIEKGYREKGQAIEYSGLTETQLRAA
                                                                                                                       373
                                                                                                                                                     391 VEAYNDFEEFSVLTTQNYAKFYYAKNKNGSSQVVYCFNADLKSPPDSEDGGKTMTPDFTT 450
                                                                                                                                                                                        315 IGLITYMRTDSTRISDTAKAEAKQYITDKY--GESYTSKRKASGKQGDQDAHEAIRPSST
                                                                                                                                                                                                                            343
                                                                                                                                                                                                                                                                                        286 NQPQTTSVLIRKYAIGDYSKLLEGATLQLTGDNVNSFQAR--VFSSNDIGERIEL-SDGT
                                                                                                                                                                                                                                                                                                                             213 WTIEGEFRYKKSKFNAKFLHYKNKPFKLKTKKDVEKITAALDGDQFEITNVTKKEKTRNP
                                                                                                                                                                                                                                                                                                                                                             245
                                                                                                                                                                                                                                                                                                                                                                                               156 DAQQ----ARRILDRLVGYNISPVLWKKVKKGLSAGRVQSVALRLVIDRENEIRNFKPEEY
                                                                                                                                                                                                                                                                                                                                                                                                                               213 STSQLSLMRQALKQLIDPNLATKMPKQV------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               170 IMEGLEPLNAIRVTQEAVWYYS------DNAPISNP-----DESFK--RESESNLV 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   112 KKAFPLGSDSSVKKWYKKHDGISTK--FEDYAMSPRITGDELNQKLRAVMYNGHPQNANG 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Gimmi, Edward R
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 610-270-5090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                       MRTPDDMKSF - - LTKDQYRLYKLIWERFVASQMA - -
                                                                                                                                                                                                                      YTL-----TELNSPAGYSIAEPITFKVEAGKVYTI---IDGKQIENPNKEIVEPYS-
                                                                                                                                                                                                                                                                                                                                                             QLSIFESEDKGDKYNK---GYQN-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KKYKVIASMGHVRDLPRSQMGVDTEDNYEPKYITIRGKGPVVKELKK-----HAKKAKN 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VFLASDP----DREGEAIAWHLSKILELEDSKENRVVFNEITKDAVKESFKNPREIEMNLV
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King of Prussia
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19.8%;
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                                                                                                                                                                                                                                                          -----TLQQEAARKLNFKARKTMMVAQQLYEGIDLKKQGT
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Pred. No. 0.069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                            ---LLSGGLVPTKPPTPGDPPMPP
                                                   ---FRANGQTIKFKGF------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      228;
                                                                                                                       ---PAILD---TVSLDITQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels 221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 711;
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                                                                                     510
                   563
                                                   434
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; TOPOLOGY: linear
; MOLECULE TYPE: prof
US-08-259-264-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/08259264
Patent No. 5650293
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (617) 227-59 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (617) 227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: pp60PIK: A DOWNSTREAM NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: ASCII text CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                      271 VPTKPPTPGDPPMPPNQPQT-TSVLIRKYAIGDYSKLLEGATLQLTGDNVNSFQARVFSS 329
416 NKNGSSQVVYCFNADLKSPPDSEDGGKTMTPDFTTGEVKYTHIAGRDLFKYTVKPRDTDP 475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      596 -GEDCEVCGSP 605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                543 VEFTVNMETLLDKIAEGDITWRKVIDGFFSSFKQDV--ERAEEEMEKIEIKDEPA----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         435 --MTLYVETKDDSDSEKENKLPKLEQ----GD-----KVTATQIEPAQHYTQPPPRYTE 482
                                    140 QYNPKLDVKLTYPVSRFQQDQLVKEDNIDAVGKNLQEFHSQYQEKS----KEYDRLY--E 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Myers, Louis
REGISTRATION NUMBER: 35
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                               30 LPPKPPKPMTPAVTNGMKDSFISLQDAEWYWGDISR-----EEVNDKLRDMPDGTFLV 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 10-JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                             ----KVEAGKVYTIIDGKQ---IENPNKEIVEPYSVEAYNDFEEFSVLTTQNYAKFYYAK 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ARLVKTLEELKIGRPSTYAPTIDTIQKRNYVKLESKRFVPTELGEIVHEQVKEYFPEIID 542
                                                                                                                    RDASTKMQ----GDYTLTLRKGGNNKLIKIYHRDGKYGFSEPLTFTSVVELINHYHHESLA 139
                                                                                                                                                         NDIGERIELSDGTYTLT------------------ 362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HGESLTLQGLP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BOSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MASSACHUSETTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                      454 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: LAHIVE & COCKFIELD 60 STATE STREET, SUITI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (617)
                                                                                                                                                                                                                                                                             Conservative
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                             protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                227-5941
                                                                                                                                                                                                                                                                           3.1%; Score 123; DB 1; Length 454: 19.4%; Pred. No. 0.054; ative 78; Mismatches 162; Indels 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/08/259, 264
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                                                                                                                                                                                                                                                                               162; Indels 142;
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Search completed: June 7, 2001, 00:17:32 Job time: 2866 sec

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